

SK



PCT09

## RAW SEQUENCE LISTING

DATE: 01/24/2002

PATENT APPLICATION: US/09/674,266A

TIME: 13:44:30

Input Set : D:\Albre3.app

Output Set: N:\CRF3\01242002\I674266A.raw

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3 <110> APPLICANT: SPECHT, THOMAS
4     HINZMANN, BERND
5     SCHMITT, ARMIN
6     PILARSKY, CHRISTIAN
7     DAHL, EDGAR
8     ROSENTHAL, ANDRE
10 <120> TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM PANCREAS TUMOR TISSUE
12 <130> FILE REFERENCE: ALBRE 3
14 <140> CURRENT APPLICATION NUMBER: 09/674,266A
15 <141> CURRENT FILING DATE: 2000-10-30
17 <160> NUMBER OF SEQ ID NOS: 661
19 <170> SOFTWARE: PatentIn Ver. 2.1
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23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
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53 <213> ORGANISM: Homo sapiens
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63 atgaaaattg tggctcctact acctgtgtt ttgttggcaa catttccgag aaagcttcag 480
64 acatgcttat aagacaactc ttagctaaat gtggtttggt tttgagctgg aagagagtac 540
65 aagggtgcttc cggaaagctt caagccttcg gattctgtga gtacaaggag ccagaatcta 600
66 ccctccgtgc actcagatta ttaçatgacc tgcaaattgg agagaaaaag ctactcgta 660
67 aagttgatgc aaagacaaag gcacagctgg atgaatggaa agcaaagaag aaagcttcta 720
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69 caaagaggag agatcagatg attaaagggg ctattgaagt tttaattcgt gaatactcca 840
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72 actaaggagg atataaatgc tatagaaatg gaagaagaca aaagagacct gatattctga 1020
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79 <213> ORGANISM: Homo sapiens
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110 &lt;211&gt; LENGTH: 2331

111 &lt;212&gt; TYPE: DNA

112 &lt;213&gt; ORGANISM: Homo sapiens

114 &lt;400&gt; SEQUENCE: 4

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117 atggaggata aaacagagaa atggtcttcc cagcctgaag gtccacttaa attgaaagct 180
118 tcaagtactg atatgccatc ccagatttct gtggttaatg tggatcaact gtgggaagat 240
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120 agctcagagg atgatgtgtt catccccact gtgaggggaa tgcatgtcc agaggccaat 360
121 attgatacag ccctttgtaa ggaaagtccg gggctctggg gagccagcat cctgaaggca 420
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123 atttcaaagg tcagagtgc ttttcagggt gtcagggtt aaagtcaaga ggtcactata 540
124 cacagcatag tgacaccaga gttttagat ctctcagtac ccaggacttt ttccactcag 600
125 attgtgcggg aatcagagat cccacgtca gagattcaaa caccttcgta cggattttcc 660
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152 atgtacacgc tgcagccaga atgcagatgg agctggcttg gctgttccct ggatgggcaa 2280
153 taaagaaagt gctgcatccc aaaaaaaaaa aaaaagtaaa aaaaaaagg g 2331

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156 &lt;210&gt; SEQ ID NO: 5

157 &lt;211&gt; LENGTH: 1925

158 &lt;212&gt; TYPE: DNA

159 &lt;213&gt; ORGANISM: Homo sapiens

161 &lt;400&gt; SEQUENCE: 5

162 aataaaaaaa attgtattta cttagaagca ttcagaatgt caacaaaaca gccgcaattt 60

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165 aaaaaaaaaa acaaaaaaca aaactaccat ccccatatat aactaatttg tgctgtgcac 240
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167 ggtagtggtg tattgaaaat accaccagga cagggtatc taaagacaca ttcggtagt 360
168 tgttaactat acaaaaaaag acaactgtaca gtttaaaaaa aaatcttaca cagccttaca 420
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192 cttaaacacgt ccggtctgaa gtttctccga gtaaacaaag atgagggaca aaagccactc 1860
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199 &lt;212&gt; TYPE: DNA

200 &lt;213&gt; ORGANISM: Homo sapiens

202 &lt;400&gt; SEQUENCE: 6

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205 aagtgtcggc aagagcgatg tctactactt cagtccaagt ggtaagaagt tcagaagcaa 180
206 gcctcagttg gcaaggtaac tgggaaatac tgttgatctc agcagttttg acttcagaac 240
207 tggaaagatg atgcctagta aattacagaa gaacaaacag agactgcgaa acgatcctct 300
208 caatcaaaat aagggttaac cagacttgaa tacaacattg ccaattagac aaacagcatc 360
209 aatttttcaa caaccggtta ccaaagtcac aaatcatcct agtaataaag tgaaatcaga 420
210 cccacaacga atgaatgaac agccacgtca gcttttctgg gagaagaggc tacaaggact 480
211 tagtgcatac gatgtaacag aacaaattat aaaaaccatg gaactacca aaggtcttca 540
212 aggagttggc ccaggtagca atgatgagac ctttttatct gctgttgcca gtgctttgca 600
213 cacaagctct gcgccaatca cagggaagt ctccgctgct gtggaaaaga accctgctgt 660
214 ttggcttaac acatctcaac cctctgcaa agcttttatt gtcacagatg aagacatcag 720

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217 agcctaagaa tatgatcagg taacttttga ccgactttcc ccaagagaaa attcctagaa 900
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221 tcaagcagga ccctaagatg aagctgagct tttgatgcca ggtgcaatct actggaaatg 1140
222 tagcacttac gtaaaacatt tgtttccccc acagttttaa taagaacaga tcaggaattc 1200
223 taaataaatt tcccagttaa agattattgt gacttactg tatataaaca tttttttata 1260
224 ctttattgaa aggggacacc tgtacattct tccatcatca ctgtaaagac aaataaatga 1320
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231 <213> ORGANISM: Homo sapiens
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236 cacagccagc agcctttcca cagtcactgc ccttcccgcg gtccccagcc ttccctacgg 180
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244 <210> SEQ ID NO: 8
245 <211> LENGTH: 1020
246 <212> TYPE: DNA
247 <213> ORGANISM: Homo sapiens
249 <400> SEQUENCE: 8
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252 gcagggtggg gaggaatcc agagttgcca tggagaaaat tccagtgtca gcattcttg 180
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259 ttgacctatc tctgacagtt agagccgata tcaactggaag atattcaaac cgtctctatg 600
260 cttacgaacc tgcagatata gctctgttgc ttgacaacat gaagaaagct ctcaagttgc 660
261 tgaagactga attgtaaaga aaaaaaatct ccaagccctt ctgtctgtca ggccttgaga 720
262 cttgaaacca gaagaagtgt gagaagactg gctagtgtgg aagcatagtg aacacactga 780
263 ttaggttatg gtttaattgt acaacaacta tttttaaga aaaacaagtt ttagaaattt 840
264 ggtttcaagt gtacatgtgt gaaaacaata ttgtatacta ccatagtgag ccatgatttt 900
265 ctaaaaaaaaa aaataaatgt tttgggggtg ttctgttttc tcaaaaaaaaa aaaaaaaaaa 960
266 aaaaaaaaaa aaaaaaaaaa aaaaattgcc cccaagggga cgggttacaa ttggggggcg 1020
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270 <211> LENGTH: 718
271 <212> TYPE: DNA

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Use of n and/or Xaa has been detected in the Sequence Listing.  
 Review the Sequence Listing to insure a corresponding  
 explanation is presented in the <220> to <223> fields of  
 each sequence using n or Xaa.

## VERIFICATION SUMMARY

DATE: 01/24/2002

PATENT APPLICATION: US/09/674,266A

TIME: 13:44:32

Input Set : D:\Albre3.app

Output Set: N:\CRF3\01242002\I674266A.raw

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L:2997 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (97) SEQUENCE:  
L:3680 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (121) SEQUENCE:  
L:3686 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (122) SEQUENCE:  
L:4047 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (141) SEQUENCE:  
L:4268 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (150) SEQUENCE:  
L:4331 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (152) SEQUENCE:  
L:11912 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (342) SEQUENCE:  
L:11918 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (343) SEQUENCE:  
L:11924 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (344) SEQUENCE:  
L:14751 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (423) SEQUENCE:  
L:14757 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (424) SEQUENCE:  
L:14763 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (425) SEQUENCE:  
L:15730 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (448) SEQUENCE:  
L:15736 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (449) SEQUENCE:  
L:15742 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (450) SEQUENCE:  
L:18377 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (522) SEQUENCE:  
L:18383 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (523) SEQUENCE:  
L:18389 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (524) SEQUENCE:  
L:18395 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (525) SEQUENCE:  
L:18401 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (526) SEQUENCE:  
L:18407 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (527) SEQUENCE:  
L:19882 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (585) SEQUENCE:  
L:19888 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (586) SEQUENCE:  
L:19894 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (587) SEQUENCE:  
L:20253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:600  
L:20254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:600  
L:20255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:600  
L:20258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:600  
L:20675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20676 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20680 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:607  
L:20952 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20954 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20957 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20958 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20959 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:20962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:609  
L:21049 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21052 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/674,266A

DATE: 01/24/2002

TIME: 13:44:32

Input Set : D:\Albre3.app

Output Set: N:\CRF3\01242002\I674266A.raw

L:21062 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21068 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21069 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21070 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21071 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21072 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21073 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21074 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:610  
L:21449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21455 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614  
L:21484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:614